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Figure 1

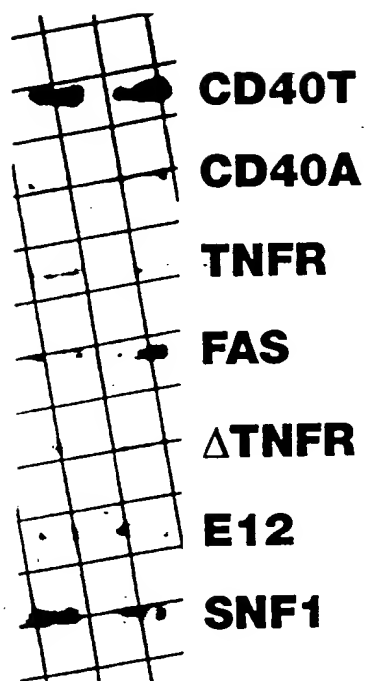


Figure 2

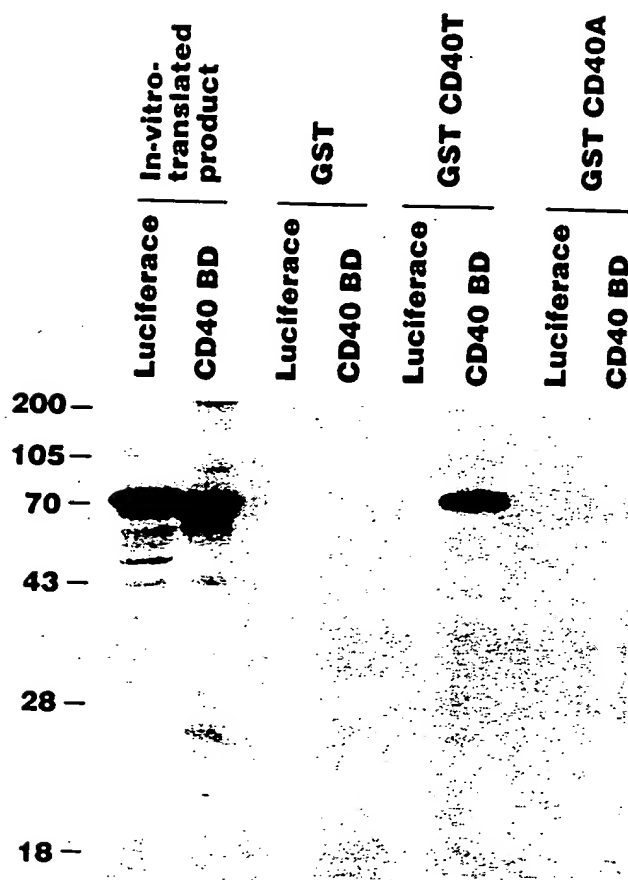


Figure 3A

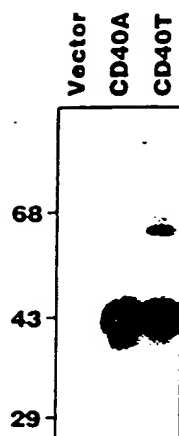


Figure 3B

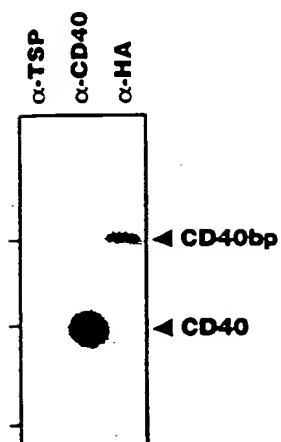


Figure 3C

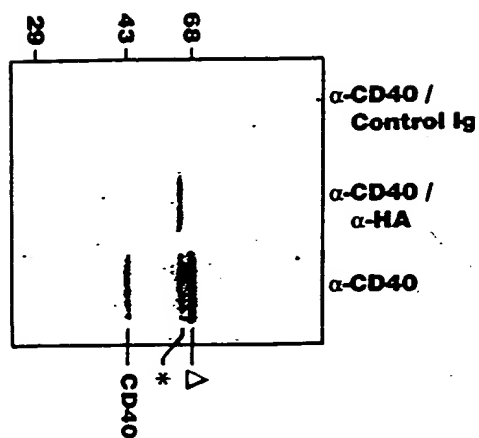


Figure 3D

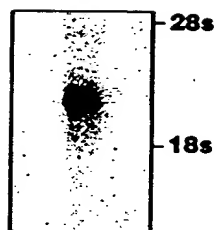
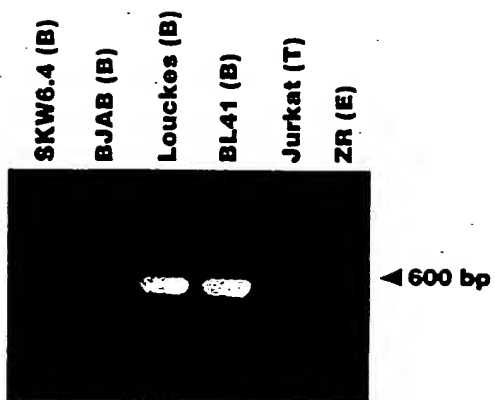


Figure 3E



MESSKKMDS PGALQTNPPLKLHTDRSAGTPVFVPEQGGYKEKFKTVEDKYKCEK 55  
 CHLVLCSPKQTECGHREFCESCMAALLSSSSPKCTACQESIIVKDKVFKDNQCKREI 110  
 LALQIYCRNESRGCAEQMLGLHVLHLLKNDCHFEELPCVVRPDCKEKVLRKDLRDHV 165  
 EKAÇKYREATÇSHÇKSQVPMIALQKHEDTDÇPÇVVVSÇPHKÇSVQTLRLRSELSAH 220  
 LSEÇVNAPSTÇSFKRYGÇVFQGTNQIQIKAHEASSAVQHVNLLKEWSNLEKKVSL 275  
 LQNESVEKNKSIQSLHNQICSEFEIEIERQKEMLRNNESEKILHLQRVIDSQAEEKLK 330  
 ELDKERIRPFQNWEEADSMKSSVESLQNRVTELESVDKSAQVARN TGLESQLS 385  
 RHDQMLSVHDIRLADMDLGFQVLETA SYNGVLIWKIRDYKRRKQEA VMGKTL SLY 440  
 SQPFYTG YFGYKMCARVYLN GDMGKGT HLSLFFVIMRGEYDALLPWPFKQKVT L 495  
 MLMDQGSRRRHLGD AFKPD PNSSSFKKPTGEMN IASGCPVFVAQTVLENGTYIKD 550  
 DTIFIKVIVDTS DLPDP 567

Figure 4A

49	DKYKCEKCHLVLCSPKQT-EC--GHRFCESCMALLSSSSPKCTACQ-ESIVK	97	CD40bp
30	AKYLCSACKNILRRPFQA-QC--GHRVCSFCLTSILSSGPQNCAACVYEGLYE	79	TRAF2
289	KSISCQICEHILADPVET-NC--KHVFCRVCI LRCLKVMGSYCPSCR-YPCFP	337	RAG1
15	SELMCPICLDMLKNTMTTKEC--LHRFCSDCIVTALRSGNKECPTCR-KKLVS	64	RING1
12	EEVTCPICLDPPFVEPVS I-EC--GHSFCQECISQVGKGGSVCPVCR-QRFL L	60	52kd RNP
30	QAFRCHVCKDFYDSPMLT-SC--NHTFCSLCIRRCLSV-DSKCP LCR-ATDQE	77	UVS-2
23	NKYTCPICFEFIYKKQIY-QCKSGHHACKECWEKSLET-KKECMTCK-SVVNS	72	DG17

Figure 4B

Figure 4C

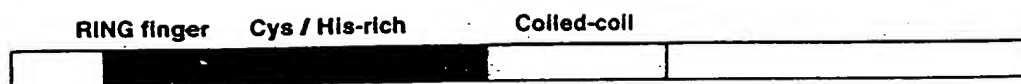
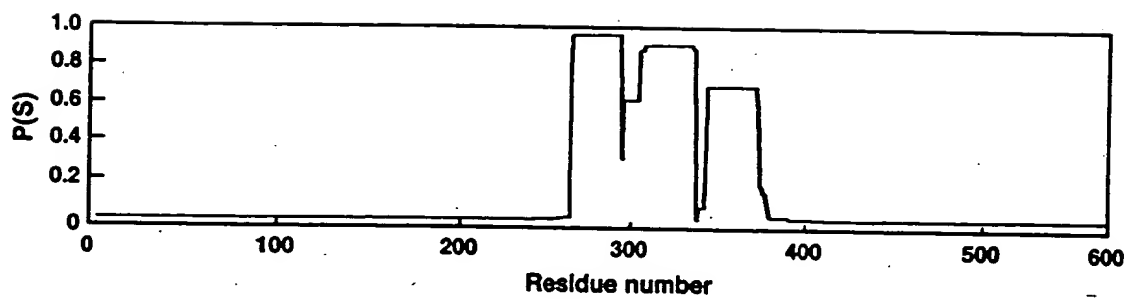


Figure 4D



[illegible]

**Figure.5**

$$=$$